
Sequence Listing was accepted.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866)

217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: [year=2008; month=5; day=7; hr=19; min=32; sec=44; ms=104;]

Validated By CRFValidator v 1.0.3

Application No: 10809689 Version No: 2.0

Input Set:

Output Set:

Started: 2008-04-23 17:58:38.189

Finished: 2008-04-23 17:58:44.403

Elapsed: 0 hr(s) 0 min(s) 6 sec(s) 214 ms

Total Warnings: 38

Total Errors: 10

No. of SeqIDs Defined: 162

Actual SeqID Count: 162

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E	341	'Xaa' position not defined SEQID (22) POS (16)									
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Input Set:

Output Set:

Started: 2008-04-23 17:58:38.189 **Finished:** 2008-04-23 17:58:44.403

Elapsed: 0 hr(s) 0 min(s) 6 sec(s) 214 ms

Total Warnings: 38

Total Errors: 10

No. of SeqIDs Defined: 162

Actual SeqID Count: 162

Error code Error Description

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<150>	US 09/610,134
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Tyr Val Glu Asn Gly Leu Ile Ser Arg Val Leu Asp Gly Leu Val Met 50 55 60

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Phe Asn Lys Ser Tyr Ala Thr Phe Glu Asp Glu Glu Ala Ala Arg Lys 40

Asn Phe Leu Glu Ser Val Lys Tyr Val Gln Ser Asn Gly Gly Ala Ile

Asn His Leu Ser Asp Leu Ser Leu Asp Glu Phe Lys Asn Arg Phe Leu 70 75

Met Ser Ala Glu Ala Phe Glu His Leu Lys Thr Gln Phe Asp Leu Asn

Ala Glu Thr Asn Ala Cys Ser Ile Asn Gly Asn Ala Pro Ala Glu Ile 105 100

Asp Leu Arg Gln Met Arg Thr Val Thr Pro Ile Arg Met Gln Gly Gly 125

115 120

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Tyr 145	Leu	Ala	Tyr	Arg	Asn 150	Gln	Ser	Leu	Asp	Leu 155	Ala	Glu	Gln	Glu	Leu 160
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Arg	Tyr	Val 195	Ala	Arg	Glu	Gln	Ser 200	Суз	Arg	Arg	Pro	Asn 205	Ala	Gln	Arg
Phe	Gly 210	Ile	Ser	Asn	Tyr	Cys 215	Gln	Ile	Tyr	Pro	Pro 220	Asn	Val	Asn	Lys
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Gly	Ile	Lys	Asp	Leu 245	Asp	Ala	Phe	Arg	His 250	Tyr	Asp	Gly	Arg	Thr 255	Ile
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Arg	Asp	Gln	Val 20	Asp	Val	Lys	Asp	Суз 25	Ala	Asn	His	Glu	Ile 30	Lys	Lys

Val Leu Val Pro Gly Cys His Gly Ser Glu Pro Cys Ile Ile His Arg

Gly Lys Pro Phe Gln Leu Glu Ala Val Phe Glu Ala Asn Gln Asn Thr 50 Lys Thr Ala Lys Ile Glu Ile Lys Ala Ser Ile Asp Gly Leu Glu Val 70 75 Asp Val Pro Gly Ile Asp Pro Asn Ala Cys His Tyr Met Lys Cys Pro 90 Leu Val Lys Gly Gln Gln Tyr Asp Ile Lys Tyr Thr Trp Asn Val Pro 100 105 Lys Ile Ala Pro Lys Ser Glu Asn Val Val Val Thr Val Lys Val Met 120 Gly Asp Asp Gly Val Leu Ala Cys Ala Ile Ala Thr His Ala Lys Ile 135 Arg Asp 145 <210> 21 <211> 261 <212> PRT <213> Dermatophagoides pteronyssinus <400> 21 Met Ile Ile Tyr Asn Ile Leu Ile Val Leu Leu Ala Ile Asn Thr 5 Leu Ala Asn Pro Ile Leu Pro Ala Ser Pro Asn Ala Thr Ile Val Gly 20 25 Gly Glu Lys Ala Leu Ala Gly Glu Cys Pro Tyr Gln Ile Ser Leu Gln 40 Ser Ser Ser His Phe Cys Gly Gly Thr Ile Leu Asp Glu Tyr Trp Ile Leu Thr Ala Ala His Cys Val Ala Gly Gln Thr Ala Ser Lys Leu Ser 70 75 Ile Arg Tyr Asn Ser Leu Lys His Ser Leu Gly Glu Lys Ile Ser Val Ala Lys Ile Phe Ala His Glu Lys Tyr Asp Ser Tyr Gln Ile Asp 100 105

Asn Asp Ile Ala Leu Ile Lys Leu Lys Ser Pro Met Lys Leu Asn Gln

125

120

115

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Leu Leu Met Glu Arg Ile His Glu Gln Ile Lys Lys Gly Glu Leu Ala 35 40 45

Leu Phe Tyr Leu Gln Glu Gln Ile Asn His Phe Glu Glu Lys Pro Thr 50 55 60

Lys Glu Met Lys Asp Lys Ile Val Ala Glu Met Asp Thr Ile Ile Ala 65 70 75 80

Met Ile Asp Gly Val Arg Gly Val Leu Asp Arg Leu Met Gln Arg Lys 85 90 95

Asp Leu Asp Ile Phe Glu Gln Tyr Asn Leu Glu Met Ala Lys Lys Ser 100 105 110

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L 5 10 15

Ser Leu Met Lys 20

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Ala Asp Pro Ile His Tyr Asp Lys Ile Thr Glu Glu Ile Asn Lys Ala 20 25 30

Val Asp Glu Ala Val Ala Ala Ile Glu Lys Ser Glu Thr Phe Asp Pro 35 40 45

Met Lys Val Pro Asp His Ser Asp Lys Phe Glu Arg His Ile Gly Ile 50 55 60

Ile Asp Leu Lys Gly Glu Leu Asp Met Arg Asn Ile Gln Val Arg Gly 65 70 75 80

Leu Lys Gln Met Lys Arg Val Gly Asp Ala Asn Val Lys Ser Glu Asp 85 90 95

Gly Val Val Lys Ala His Leu Leu Val Gly Val His Asp Asp Val Val
100 105 110

Ser Met Glu Tyr Asp Leu Ala Tyr Lys Leu Gly Asp Leu His Pro Asn 115 120 125

Thr His Val Ile Ser Asp Ile Gln Asp Phe Val Val Glu Leu Ser Leu 130 135 140

Glu Val Ser Glu Glu Gly Asn Met Thr Leu Thr Ser Phe Glu Val Arg 145 150 155 160

Gln Phe Ala Asn Val Val Asn His Ile Gly Gly Leu Ser Ile Leu Asp 165 170 175

Pro Ile Phe Ala Val Leu Ser Asp Val Leu Thr Ala Ile Phe Gln Asp 180 185 190

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Glu Leu Glu Arg Asn Asn Gln

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Lys Asn Tyr Ala Thr Val Glu Glu Glu Val Ala Arg Lys Asn Phe 35 40 45

Leu Glu Ser Leu Lys Tyr Val Glu Ala Asn Lys Gly Ala Ile Asn His 50 60

Leu Ser Asp Leu Ser Leu Asp Glu Phe Lys Asn Arg Tyr Leu Met Ser 65 70 75 80

Ala Glu Ala Phe Glu Gln Leu Lys Thr Gln Phe Asp Leu Asn Ala Glu 85 90 95

Thr Ser Ala Cys Arg Ile Asn Ser Val Asn Val Pro Ser Glu Leu Asp 100 105 110

Leu Arg Ser Leu Arg Thr Val Thr Pro Ile Arg Met Gln Gly Gly Cys 115 120 125

Gly Ser Cys Trp Ala Phe Ser Gly Val Ala Ala Thr Glu Ser Ala Tyr

130 135 140

Leu Ala Tyr Arg Asn Thr Ser Leu Asp Leu Ser Glu Gln Glu Leu Val
145 150 150 155 160

Asp Cys Ala Ser Gln His Gly Cys His Gly Asp Thr Ile Pro Arg Gly \$165\$ \$170\$ \$175\$

Ile Glu Tyr Ile Gln Gln Asn Gly Val Val Glu Glu Arg Ser Tyr Pro\$180\$ \$190\$

Tyr Val Ala Arg Glu Gln Arg Cys Arg Arg Pro Asn Ser Gln His Tyr 195 200 205

Gly Ile Ser Asn Tyr Cys Gln Ile Tyr Pro Pro Asp Val Lys Gln Ile 210 215 220

Arg Glu Ala Leu Thr Gln Thr His Thr Ala Ile Ala Val Ile Ile Gly 225 230 235 240

Ile Lys Asp Leu Arg Ala Phe Gl